



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/748,128
Source: FWO
Date Processed by STIC: 1/13/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM. ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efc/efs/downloads/documents.htm>>, EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/748,128

DATE: 01/13/2004

TIME: 10:09:55

Input Set : A:\66821-058.TXT

Output Set: N:\CRF4\01092004\J748128.raw

4 <110> APPLICANT: Reed, John C.
 5 Houghten, Richard A.
 6 Nefzi, Adel
 7 Ostresh, John M.
 8 Pinilla, Clemencia
 9 Welsh, Kate
 11 <120> TITLE OF INVENTION: Methods and Compositions for
 12 Derepression of IAP-Inhibited Caspase
 15 <130> FILE REFERENCE: 66821-058
 C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/748,128
 C--> 17 <141> CURRENT FILING DATE: 2003-12-24
 17 <150> PRIOR APPLICATION NUMBER: US 60/331,957
 18 <151> PRIOR FILING DATE: 2001-11-21
 20 <150> PRIOR APPLICATION NUMBER: US 10/302,811
 21 <151> PRIOR FILING DATE: 2002-11-21
 23 <160> NUMBER OF SEQ ID NOS: 16
 25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 5
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Artificial Sequence
 32 <220> FEATURE:
 33 <223> OTHER INFORMATION: synthetic construct
 W--> 35 <221> NAME/KEY: VARIANT
 36 <222> LOCATION: 4
 37 <223> OTHER INFORMATION: Xaa = Any Amino Acid
 W--> 39 <400> 1
 W--> 40 Gln Ala Cys Xaa Gly
 41 1 5
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 4
 46 <212> TYPE: PRT
 47 <213> ORGANISM: Artificial Sequence
 49 <220> FEATURE:
 50 <223> OTHER INFORMATION: synthetic construct
 52 <400> SEQUENCE: 2
 53 Asp Glu Val Asp
 54 1
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 4
 59 <212> TYPE: PRT
 60 <213> ORGANISM: Artificial Sequence
 62 <220> FEATURE:

Does Not Comply
Corrected Diskette Needed

P.S

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Input Set : A:\66821-058.TXT

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63 <223> OTHER INFORMATION: synthetic construct
65 <400> SEQUENCE: 3
66 Tyr Val Ala Asp
67 1
70 <210> SEQ ID NO: 4
71 <211> LENGTH: 4
72 <212> TYPE: PRT
73 <213> ORGANISM: Artificial Sequence
75 <220> FEATURE:
76 <223> OTHER INFORMATION: synthetic construct
78 <400> SEQUENCE: 4
79 Ala Val Pro Ile
80 1
83 <210> SEQ ID NO: 5
84 <211> LENGTH: 7
85 <212> TYPE: PRT
86 <213> ORGANISM: Artificial Sequence
88 <220> FEATURE:
89 <223> OTHER INFORMATION: synthetic construct
W--> 91 <221> NAME/KEY: AMIDATION
92 <222> LOCATION: 7
93 <223> OTHER INFORMATION: at the C-terminus
W--> 95 <221> MOD_RES
96 <222> LOCATION: 1
97 <223> OTHER INFORMATION: hydrogenated at the N-terminus
W--> 99 <400> 5
100 Ala Val Pro Ile Ala Gln Lys
101 1 5
104 <210> SEQ ID NO: 6
105 <211> LENGTH: 4
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: synthetic peptide
112 <400> SEQUENCE: 6
113 Ala Val Pro Ser
114 1
117 <210> SEQ ID NO: 7
118 <211> LENGTH: 6
119 <212> TYPE: PRT
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: synthetic peptide
W--> 125 <221> NAME/KEY: VARIANT
126 <222> LOCATION: (1)...(2)
127 <223> OTHER INFORMATION: Xaa = any amino acid
W--> 129 <400> 7
W--> 130 Xaa Xaa Ala Ala Trp Trp
131 1 5

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Input Set : A:\66821-058.TXT

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134 <210> SEQ ID NO: 8
135 <211> LENGTH: 6
136 <212> TYPE: PRT
137 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: synthetic peptide
W--> 142 <221> NAME/KEY: VARIANT
143 <222> LOCATION: (1)...(2)
144 <223> OTHER INFORMATION: Xaa = any amino acid
W--> 146 <400> 8
W--> 147 Xaa Xaa Gly Ala Trp Trp
148 1 5
151 <210> SEQ ID NO: 9
152 <211> LENGTH: 6
153 <212> TYPE: PRT
154 <213> ORGANISM: Artificial Sequence
156 <220> FEATURE:
157 <223> OTHER INFORMATION: synthetic peptide
W--> 159 <221> NAME/KEY: VARIANT
160 <222> LOCATION: (1)...(2)
161 <223> OTHER INFORMATION: Xaa = any amino acid
W--> 163 <400> 9
W--> 164 Xaa Xaa Arg Ala Trp Trp
165 1 5
168 <210> SEQ ID NO: 10
169 <211> LENGTH: 6
170 <212> TYPE: PRT
171 <213> ORGANISM: Artificial Sequence
173 <220> FEATURE:
174 <223> OTHER INFORMATION: synthetic peptide
W--> 176 <221> NAME/KEY: VARIANT
177 <222> LOCATION: (1)...(2)
178 <223> OTHER INFORMATION: Xaa = any amino acid
W--> 180 <400> 10
W--> 181 Xaa Xaa Cys Lys Trp Trp
182 1 5
185 <210> SEQ ID NO: 11
186 <211> LENGTH: 6
187 <212> TYPE: PRT
188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: synthetic peptide
W--> 193 <221> NAME/KEY: VARIANT
194 <222> LOCATION: 1, 2
195 <223> OTHER INFORMATION: Xaa = Any Amino Acid
W--> 197 <400> 11
W--> 198 Xaa Xaa Phe Trp Trp Trp
199 1 5
202 <210> SEQ ID NO: 12

RAW SEQUENCE LISTING

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Input Set : A:\66821-058.TXT

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203 <211> LENGTH: 6
204 <212> TYPE: PRT
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: synthetic peptide
W--> 210 <221> NAME/KEY: VARIANT
211 <222> LOCATION: 1, 2
212 <223> OTHER INFORMATION: Xaa = Any Amino Acid
W--> 214 <400> 12
W--> 215 Xaa Xaa Leu Trp Trp Trp
216 1 5
219 <210> SEQ ID NO: 13
220 <211> LENGTH: 6
221 <212> TYPE: PRT
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: synthetic peptide
W--> 227 <221> NAME/KEY: VARIANT
228 <222> LOCATION: 1, 2
229 <223> OTHER INFORMATION: Xaa = Any Amino Acid
W--> 231 <400> 13
W--> 232 Xaa Xaa Trp Leu Trp Trp
233 1 5
236 <210> SEQ ID NO: 14
237 <211> LENGTH: 6
238 <212> TYPE: PRT
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: synthetic peptide
W--> 244 <221> NAME/KEY: VARIANT
245 <222> LOCATION: 1, 2
246 <223> OTHER INFORMATION: Xaa = Any Amino Acid
W--> 248 <400> 14
W--> 249 Xaa Xaa Trp Trp Trp Trp
250 1 5
253 <210> SEQ ID NO: 15
254 <211> LENGTH: 6
255 <212> TYPE: PRT
256 <213> ORGANISM: Artificial Sequence
258 <220> FEATURE:
259 <223> OTHER INFORMATION: synthetic peptide
W--> 261 <221> NAME/KEY: VARIANT
262 <222> LOCATION: (1)...(2)
263 <223> OTHER INFORMATION: Xaa=any amino acid
W--> 265 <400> 15
W--> 266 Xaa Xaa Leu Lys Trp Trp
267 1 5
270 <210> SEQ ID NO: 16
271 <211> LENGTH: 68

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Input Set : A:\66821-058.TXT

Output Set: N:\CRF4\01092004\J748128.raw

272 <212> TYPE: PRT
 273 <213> ORGANISM: Artificial Sequence
 275 <220> FEATURE:
 276 <223> OTHER INFORMATION: consensus sequence
 W--> 278 <221> NAME/KEY: VARIANT
 279 <222> LOCATION: 1, 3, 6, 8, 10, 11, 14, 16, 18, 21, 24, 30, 33, 35, 37, 40, 42=44, 46, 47, (see below)
 280 49=51, 53=57, 59, 61, 62, 64, 66
 281 <223> OTHER INFORMATION: Xaa=Any amino acid
 W--> 283 <221> VARIANT
 284 <222> LOCATION: 15
 285 <223> OTHER INFORMATION: Xaa=any amino acid that may or may not be present
 W--> 287 <221> VARIANT
 288 <222> LOCATION: 3, 17, 28, 29, 45, 68
 289 <223> OTHER INFORMATION: Xaa=a hydrophobic amino acid
 W--> 291 <221> VARIANT
 292 <222> LOCATION: 7
 293 <223> OTHER INFORMATION: Xaa=serine or threonine
 W--> 295 <221> VARIANT
 296 <222> LOCATION: 8, 67
 297 <223> OTHER INFORMATION: Xaa=phenylalanine or tyrosine
 W--> 299 <221> VARIANT
 300 <222> LOCATION: (12)...(12)
 301 <223> OTHER INFORMATION: Xaa=proline that may or may not be present
 W--> 303 <221> VARIANT
 304 <222> LOCATION: (52)...(52)
 305 <223> OTHER INFORMATION: Xaa=aspartic or glutamic acid
 W--> 307 <221> VARIANT
 308 <222> LOCATION: (66)...(66)
 309 <223> OTHER INFORMATION: Xaa=a basic amino acid
 W--> 311 <400> 16
 W--> 312 Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Xaa Xaa
 313 1 5 10 15
 314 Xaa Xaa Xaa Xaa Xaa Leu Ala Xaa Ala Gly Phe Xaa Xaa Xaa Gly Xaa
 315 20 25 30
 316 Xaa Asp Xaa Val Xaa Cys Phe Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Trp
 317 35 40 45
 318 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Pro Xaa
 319 50 55
 320 Cys Xaa Xaa Xaa
 321 65

what about Xaa's at locations
 32 and 60?
 (see below)

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/748,128

DATE: 01/13/2004
 TIME: 10:09:56

Input Set : A:\66821-058.TXT
 Output Set: N:\CRF4\01092004\J748128.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 4
 Seq#:7; Xaa Pos. 1, 2
 Seq#:8; Xaa Pos. 1, 2
 Seq#:9; Xaa Pos. 1, 2
 Seq#:10; Xaa Pos. 1, 2
 Seq#:11; Xaa Pos. 1, 2
 Seq#:12; Xaa Pos. 1, 2
 Seq#:13; Xaa Pos. 1, 2
 Seq#:14; Xaa Pos. 1, 2
 Seq#:15; Xaa Pos. 1, 2
 Seq#:16; Xaa Pos. 1, 2, 3, 5, 6, 7, 8, 9, 10, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 24, 28, 29
 Seq#:16; Xaa Pos. 30, 32, 33, 35, 37, 40, 42, 43, 44, 45, 46, 47, 49, 50, 51, 52, 53, 54, 55
 Seq#:16; Xaa Pos. 56, 57, 59, 60, 61, 62, 64, 66, 67, 68

VERIFICATION SUMMARY

DATE: 01/13/2004

PATENT APPLICATION: US/10/748,128

TIME: 10:09:56

Input Set : A:\66821-058.TXT

Output Set: N:\CRF4\01092004\J748128.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:35 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:39 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:40 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:91 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:95 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:99 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:125 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:129 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:142 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:146 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:159 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:163 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:176 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:180 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:193 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:197 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:210 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:214 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:227 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:231 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13
L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:244 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:248 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:261 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:265 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:278 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:283 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:287 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:291 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:295 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:299 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:303 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:307 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:311 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:312 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
M:341 Repeated in SeqNo=16